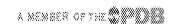
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As of Tuesday Aug 25, 2009 Mithere are 59790 Structures 🕷 | PDB Statistics 🕷

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> XrayCellQuery: cell.length_a.comparator= between cell.length_a.min= 41.22 cell.length_a.max= 41.24 cell.length b.comparator=between cell.length b.min=65.21 cell.length b.max=65.23 cell.length c.comparator=between cell.length c.min=73.37 cell.length c.max=73.39 cell.angle_alpha.comparator=between cell.angle_alpha.min=73.10 cell.angle_alpha.max=73.12 cell.angle_beta.comparator=between cell.angle_beta.min=85.57 cell.angle_beta.max=85.59 cell.angle_gamma.comparator=between cell.angle_gamma.min=85.79 cell.angle_gamma.max=85.81

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A CONSERVED PROTEIN BINDING-SITE ON BACTERIAL SLIDING CLAMPS

Characteristics

Release Date: 15-Jul-2004 Exp. Method: X-RAY DIFFRACTION

Resolution: 1.65 Å

Classification

Transferase

Compound

Molecule:

DNA POLYMERASE III

Polymer:

Type: polypeptide(L)

Chains: $\mathbb{C}(\mathbb{W})$

A. 8

2.7.7.7

Molecule:

null

Polymer: 2 Type:

polypeptide(L)

Length: 16

Length: 366

Chains:

C-TERMINUS OF DNA POLITY ELOQLI, RESIDUES 1-16 Fragment:

Other Details: C-TERMINUS (16 RESIDUES) OF DNA POLIVIE.COLL

Authors

Burnouf, D.Y., Olieric, V., Wagner, J., Fujii, S., Reinbolt, J., Fuchs,

R.P.P., Dumas, P.

[1]

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The RCSB PDB is managed by two members of the RCSB: Rutgers and UCSD, and is funded by NSF, NI GMS, DOE, NEM, NCI , NI NDS, and NI DDK.

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